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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/920,068A

DATE: 02/10/2002
TIME: 13:06:11

Input Set : A:\50125.015002.SEQLIST.TXT
Output Set: N:\CRF3\02102002\I920068A.raw

4 <110> APPLICANT: Wolf, Eckhard
5 Werner, Sabine
6 Halle, Jorn-Peter
7 Regenbogen, Johannes
8 Goppelt, Andreas
10 <120> TITLE OF INVENTION: Polypeptides or Nucleic Acids Encoding
11 These of a Family of G-Protein Coupled Receptors and their
12 Use for the Diagnosis or Treatment of Disorders, for example
13 Skin Disorders and their Use for the Identification of
14 Pharmacologically Active Substances
17 <130> FILE REFERENCE: 50125/015002
19 <140> CURRENT APPLICATION NUMBER: 09/920,068A
20 <141> CURRENT FILING DATE: 2001-08-01
22 <150> PRIOR APPLICATION NUMBER: 60/229,501
23 <151> PRIOR FILING DATE: 2000-08-31
25 <150> PRIOR APPLICATION NUMBER: DE 10038111.1
26 <151> PRIOR FILING DATE: 2000-08-04
28 <160> NUMBER OF SEQ ID NOS: 21
30 <170> SOFTWARE: FastSEQ for Windows Version 4.0
32 <210> SEQ ID NO: 1
33 <211> LENGTH: 331
34 <212> TYPE: PRT
35 <213> ORGANISM: Mus musculus
37 <400> SEQUENCE: 1
38 Met Gly Glu Ser Asn Gly Glu Ala Phe Leu Ala Phe Lys Thr Ser Ala
39 1 5 10 15
40 Ser Pro Thr Ala Pro Val Thr Thr Asn Pro Met Asp Glu Thr Leu Pro
41 20 25 30
42 Gly Ser Ile Asn Ile Arg Ile Leu Ile Pro Lys Leu Met Ile Ile Ile
43 35 40 45
44 Phe Gly Leu Val Gly Leu Met Gly Asn Ala Ile Val Phe Trp Leu Leu
45 50 55 60
46 Gly Phe His Leu Arg Arg Asn Ala Phe Ser Val Tyr Ile Leu Asn Leu
47 65 70 75 80
48 Ala Leu Ala Asp Phe Leu Phe Leu Leu Ser Ser Ile Ile Ala Ser Thr
49 85 90 95
50 Leu Phe Leu Leu Lys Val Ser Tyr Leu Ser Ile Ile Phe His Leu Cys
51 100 105 110
52 Phe Asn Thr Ile Met Met Val Val Tyr Ile Thr Gly Ile Ser Met Leu
53 115 120 125
54 Ser Ala Ile Ser Thr Glu Cys Cys Leu Ser Val Leu Cys Pro Thr Trp
55 130 135 140
56 Tyr Arg Cys His Arg Pro Val His Thr Ser Thr Val Met Cys Ala Val

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Input Set : A:\50125.015002.SEQLIST.TXT
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57 145 150 155 160
 58 Ile Trp Val Leu Ser Leu Leu Ile Cys Ile Leu Asn Ser Tyr Phe Cys
 59 165 170 175
 60 Ala Val Leu His Thr Arg Tyr Asp Asn Asp Asn Glu Cys Leu Ala Thr
 61 180 185 190
 62 Asn Ile Phe Thr Ala Ser Tyr Met Ile Phe Leu Leu Val Val Leu Cys
 63 195 200 205
 64 Leu Ser Ser Leu Ala Leu Leu Ala Arg Leu Phe Cys Gly Ala Gly Gln
 65 210 215 220
 66 Met Lys Leu Thr Arg Phe His Val Thr Ile Leu Leu Thr Leu Leu Val
 67 225 230 235 240
 68 Phe Leu Leu Cys Gly Leu Pro Phe Val Ile Tyr Cys Ile Leu Leu Phe
 69 245 250 255
 70 Lys Ile Lys Asp Asp Phe His Val Leu Asp Val Asn Leu Tyr Leu Ala
 71 260 265 270
 72 Leu Glu Val Leu Thr Ala Ile Asn Ser Cys Ala Asn Pro Ile Ile Tyr
 73 275 280 285
 74 Phe Phe Val Gly Ser Phe Arg His Gln Leu Lys His Gln Thr Leu Lys
 75 290 295 300
 76 Met Val Leu Gln Ser Ala Leu Gln Asp Thr Pro Glu Thr Ala Glu Asn
 77 305 310 315 320
 78 Met Val Glu Met Ser Ser Asn Lys Ala Glu Pro
 79 325 330
 82 <210> SEQ ID NO: 2
 83 <211> LENGTH: 321
 84 <212> TYPE: PRT
 85 <213> ORGANISM: Homo sapiens
 87 <400> SEQUENCE: 2
 88 Met Asn Gln Thr Leu Asn Ser Ser Gly Thr Val Glu Ser Ala Leu Asn
 89 1 5 10 15
 90 Tyr Ser Arg Gly Ser Thr Val His Thr Ala Tyr Leu Val Leu Ser Ser
 91 20 25 30
 92 Leu Ala Met Phe Thr Cys Leu Cys Gly Met Ala Gly Asn Ser Met Val
 93 35 40 45
 94 Ile Trp Leu Leu Gly Phe Arg Met His Arg Asn Pro Phe Cys Ile Tyr
 95 50 55 60
 96 Ile Leu Asn Leu Ala Ala Asp Leu Leu Phe Leu Phe Ser Met Ala
 97 65 70 75 80
 98 Ser Thr Leu Ser Leu Glu Thr Gln Pro Leu Val Asn Thr Thr Asp Lys
 99 85 90 95
 100 Val His Glu Leu Met Lys Arg Leu Met Tyr Phe Ala Tyr Thr Val Gly
 101 100 105 110
 102 Leu Ser Leu Leu Thr Ala Ile Ser Thr Gln Arg Cys Leu Ser Val Leu
 103 115 120 125
 104 Phe Pro Ile Trp Phe Lys Cys His Arg Pro Arg His Leu Ser Ala Trp
 105 130 135 140
 106 Val Cys Gly Leu Leu Trp Thr Leu Cys Leu Leu Met Asn Gly Leu Thr
 107 145 150 155 160
 108 Ser Ser Phe Cys Ser Lys Phe Leu Lys Phe Asn Glu Asp Arg Cys Phe

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109	165	170	175	
110	Arg Val Asp Met Val Gln Ala Ala Leu Ile Met Gly Val Leu Thr Pro			
111	180	185	190	
112	Val Met Thr Leu Ser Ser Leu Thr Leu Phe Val Trp Val Arg Arg Ser			
113	195	200	205	
114	Ser Gln Gln Trp Arg Arg Gln Pro Thr Arg Leu Phe Val Val Val Leu			
115	210	215	220	
116	Ala Ser Val Leu Val Phe Leu Ile Cys Ser Leu Pro Leu Ser Ile Tyr			
117	225	230	235	240
118	Trp Phe Val Leu Tyr Trp Leu Ser Leu Pro Pro Glu Met Gln Val Leu			
119	245	250	255	
120	Cys Phe Ser Leu Ser Arg Leu Ser Ser Ser Val Ser Ser Ser Ala Asn			
121	260	265	270	
122	Pro Val Ile Tyr Phe Leu Val Gly Ser Arg Arg Ser His Arg Leu Pro			
123	275	280	285	
124	Thr Arg Ser Leu Gly Thr Val Leu Gln Gln Ala Leu Arg Glu Glu Pro			
125	290	295	300	
126	Glu Leu Glu Gly Gly Glu Thr Pro Thr Val Gly Thr Asn Glu Met Gly			
127	305	310	315	320
128	Ala			

132	<210> SEQ ID NO: 3			
133	<211> LENGTH: 325			
134	<212> TYPE: PRT			
135	<213> ORGANISM: Mus musculus			
137	<400> SEQUENCE: 3			
138	Met Asp Ile Asp Ile Ser Ser Leu Gly Ile Tyr Ile Ile Ala Pro Asn			
139	1	5	10	15
140	Gly Ser Ser Tyr Thr Asn Ser Val Asp Cys Phe Phe Lys Ile Gln Val			
141	20	25	30	
142	Met Gly Phe Leu Ser Leu Ile Ile Ser Pro Val Gly Met Val Leu Asn			
143	35	40	45	
144	Ser Thr Val Leu Trp Phe Leu Gly Phe Gln Ile Arg Arg Asn Ala Phe			
145	50	55	60	
146	Ser Val Tyr Ile Leu Asn Leu Ala Gly Ala Asp Phe Leu Phe Leu His			
147	65	70	75	80
148	Ser Gln Phe Leu Phe Tyr Leu Leu Ala Ile Phe Pro Ser Ile Pro Ile			
149	85	90	95	
150	Gln Ile Pro Leu Phe Phe Asp Met Leu Thr Lys Phe Ala Tyr Leu Ser			
151	100	105	110	
152	Gly Leu Ser Ile Leu Ser Thr Ile Ser Ile Glu Arg Cys Leu Cys Val			
153	115	120	125	
154	Met Trp Pro Ile Trp Tyr Arg Cys Gln Arg Pro Arg His Thr Ser Ser			
155	130	135	140	
156	Val Thr Cys Ser Leu Leu Trp Ala Leu Ser Leu Leu Phe Ala Leu Leu			
157	145	150	155	160
158	Asp Gly Met Gly Cys Gly Leu Leu Phe Asn Ser Phe Asp Gln Ser Trp			
159	165	170	175	
160	Cys Leu Lys Phe Asp Leu Ile Ile Cys Ala Trp Ser Ile Val Leu Phe			
161	180	185	190	

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Input Set : A:\50125.015002.SEQLIST.TXT
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162 Val Val Leu Cys Gly Ser Ser Leu Ile Leu Leu Val Arg Ile Phe Cys
 163 195 200 205
 164 Gly Ser Gln Gln Ile Pro Val Thr Arg Leu Tyr Val Thr Ile Ala Leu
 165 210 215 220
 166 Thr Val Leu Phe Phe Leu Ile Cys Cys Leu Pro Phe Gly Ile Ser Trp
 167 225 230 235 240
 168 Ile Ile Gln Trp Ser Glu Thr Leu Ile Tyr Val Gly Phe Cys Asp Tyr
 169 245 250 255
 170 Phe His Glu Glu Leu Phe Leu Ser Cys Ile Asn Ser Cys Ala Asn Pro
 171 260 265 270
 172 Ile Ile Tyr Phe Leu Val Gly Phe Ile Arg Gln Arg Lys Phe Gln Gln
 173 275 280 285
 174 Lys Ser Leu Lys Val Leu Leu Gln Arg Ala Met Glu Asp Thr Pro Glu
 175 290 295 300
 176 Glu Glu Asn Glu Asp Met Gly Pro Ser Arg Asn Pro Glu Glu Phe Glu
 177 305 310 315 320
 178 Thr Val Cys Ser Asn
 179 325
 182 <210> SEQ ID NO: 4
 183 <211> LENGTH: 330
 184 <212> TYPE: PRT
 185 <213> ORGANISM: Homo sapiens
 187 <400> SEQUENCE: 4
 188 Met Asp Pro Thr Thr Pro Ala Trp Gly Thr Glu Ser Thr Thr Val Asn
 189 1 5 10 15
 190 Gly Asn Asp Gln Ala Leu Leu Leu Cys Gly Lys Glu Thr Leu Ile
 191 20 25 30
 192 Pro Val Phe Leu Ile Leu Phe Ile Ala Leu Val Gly Leu Val Gly Asn
 193 35 40 45
 194 Gly Phe Val Leu Trp Leu Leu Gly Phe Arg Met Arg Arg Asn Ala Phe
 195 50 55 60
 196 Ser Val Tyr Val Leu Ser Leu Ala Gly Ala Asp Phe Leu Phe Leu Cys
 197 65 70 75 80
 198 Phe Gln Ile Ile Asn Cys Leu Val Tyr Leu Ser Asn Phe Phe Cys Ser
 199 85 90 95
 200 Ile Ser Ile Asn Phe Pro Ser Phe Phe Thr Thr Val Met Thr Cys Ala
 201 100 105 110
 202 Tyr Leu Ala Gly Leu Ser Met Leu Ser Thr Val Ser Thr Glu Arg Cys
 203 115 120 125
 204 Leu Ser Val Leu Trp Pro Ile Trp Tyr Arg Cys Arg Arg Pro Arg His
 205 130 135 140
 206 Leu Ser Ala Val Val Cys Val Leu Leu Trp Ala Leu Ser Leu Leu
 207 145 150 155 160
 208 Ser Ile Leu Glu Gly Lys Phe Cys Gly Phe Leu Phe Ser Asp Gly Asp
 209 165 170 175
 210 Ser Gly Trp Cys Gln Thr Phe Asp Phe Ile Thr Ala Ala Trp Leu Ile
 211 180 185 190
 212 Phe Leu Phe Met Val Leu Cys Gly Ser Ser Leu Ala Leu Leu Val Arg
 213 195 200 205

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Input Set : A:\50125.015002.SEQLIST.TXT
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214 Ile Leu Cys Gly Ser Arg Gly Leu Pro Leu Thr Arg Leu Tyr Leu Thr
 215 210 215 220
 216 Ile Leu Leu Thr Val Leu Val Phe Leu Leu Cys Gly Leu Pro Phe Gly
 217 225 230 235 240
 218 Ile Gln Trp Phe Leu Ile Leu Trp Ile Trp Lys Asp Ser Asp Val Leu
 219 245 250 255
 220 Phe Cys His Ile His Pro Val Ser Val Val Leu Ser Ser Leu Asn Ser
 221 260 265 270
 222 Ser Ala Asn Pro Ile Ile Tyr Phe Phe Val Gly Ser Phe Arg Lys Gln
 223 275 280 285
 224 Trp Arg Leu Gln Gln Pro Ile Leu Lys Leu Ala Leu Gln Arg Ala Leu
 225 290 295 300
 226 Gln Asp Ile Ala Glu Val Asp His Ser Glu Gly Cys Phe Arg Gln Gly
 227 305 310 315 320
 228 Thr Pro Glu Met Ser Arg Ser Ser Leu Val
 229 325 330
 232 <210> SEQ ID NO: 5
 233 <211> LENGTH: 993
 234 <212> TYPE: DNA
 235 <213> ORGANISM: Mus musculus
 237 <400> SEQUENCE: 5
 238 atgggggaaa gcaatggta agcatttctt gccttaaga cctcagccctc accaacagca 60
 239 ccagtgacaa caaatccaaat ggacgaaacc ctccctggaa gtatcaacat taggattctg 120
 240 atccccaaat tgatgatcat catcttcgga ctggcggac tgatggaaa cgccattgtg 180
 241 ttctggctcc tgggcttcca cttgcgcagg aatgccttct cagtctacat cctaaacttg 240
 242 gccctggctg acttcctttt cctcctcagt agtacatag cttccacccct gtttcttctc 300
 243 aaagtttctt acctcagcat catcttcac ttgtgcttta acaccattat gatggttgtc 360
 244 tacatcacag ggataagcat gctcagtgcc atcagcactg agtgcgtcct gtctgtcctg 420
 245 tggcccaccc ggtatcgctg ccaccgtcca gtacatacat caactgtcat gtgtgctgtg 480
 246 atctgggtcc tatccctgtt gatctgcatt ctgaatagct atttctgtgc tgtcttacat 540
 247 accagatatg ataatgacaa tgagtgtctg gcaactaaca tctttaccgc ctcgtacatg 600
 248 atattttgc ttgtggctt ctgtctgtcc agcctggctc tgctggccag gttgttctgt 660
 249 ggcgctgggc agatgaagct taccagattt catgtgacca tcttgctgac cctttgggtt 720
 250 tttctccctt gcgggttgcc ctttgtcata tactgcatcc tggatttcaa gattaaggat 780
 251 gatttccatg tattagatgt taatctttt ctacgattag aagtcctgac tgctattaac 840
 252 agctgtgcca accccatcat ctacttcttc gtggcctt tcagacatca gttgaagcac 900
 253 cagaccctca aaatggttct ccagagtgca ctgcaggaca ctccctgagac agctgaaaac 960
 254 atggtagaga tgtcaagtaa caaagcagag cct 993
 256 <210> SEQ ID NO: 6
 257 <211> LENGTH: 966
 258 <212> TYPE: DNA
 259 <213> ORGANISM: Homo sapiens
 261 <400> SEQUENCE: 6
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 263 agcacagtgc acacggccata cctgggtctg agctccctgg ccatgttcac ctgcctgtgc 120
 264 gggatggcag gcaacagcat ggtgatctgg ctgtctggct ttcgaatgca caggaacccc 180
 265 ttctgcataat atatcccaa cctggcgcca gccgacccctc tcttcctt cagcatggct 240
 266 tccacgctca gcctggaaac ccagccctg gtcaatacca ctgacaaggat ccacgagctg 300
 267 atgaagagac tgatgtactt tgcctacaca gtggcctga gcctgctgac ggcacatcagc 360

Use of n and/or Xaa has been detected in the Sequence Listing.
 → Review the Sequence Listing to insure a corresponding
 explanation is presented in the <220> to <223> fields of
 each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/920,068A

DATE: 02/10/2002

TIME: 13:06:12

Input Set : A:\50125.015002.SEQLIST.TXT

Output Set: N:\CRF3\02102002\I920068A.raw

L:409 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15

L:410 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15

L:411 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15

L:412 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15